

Gp 1618

PATENT  
Docket No.: 19603/468 (CRF D-1595C)

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

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Applicants : Barany et al.

Serial No. : 08/794,851

Filed : February 4, 1997

For : DETECTION OF NUCLEIC ACID  
SEQUENCE DIFFERENCES USING THE  
LIGASE DETECTION REACTION WITH  
ADDRESSABLE ARRAYS

Examiner:  
J. Ricigliano

Art Unit:  
1618



Assistant Commissioner for Patents  
Washington, D.C. 20231

Box: Non-Fee Amendment

Sir:

Transmitted herewith is an amendment (7 pages) in the above-identified application.  
Also enclosed are:

- ☒ Correction of Correspondence Address.
- ☒ A self-addressed, prepaid postcard for acknowledging receipt.
- ☒ No additional fee is required.
- ☒ The fee has been calculated as shown below:

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TOTAL	87	MINUS	89	0	a) \$ 9= b) \$18	\$0
INDEP	1	MINUS	3	0	a) \$39= b) \$78	\$0
[ ] FIRST PRESENTATION OF MULTIPLE DEPENDENT CLAIM					see fee schedule	\$0
[ ] Extension of Time Fee (___ Months)						
TOTAL						\$0

- [ ] A check for \$\_\_\_\_\_ is enclosed to cover the above fees.
- [ ] Please charge my Deposit Account No. 14-1138 in the amount of \$\_\_\_\_\_  
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- [X] The Commissioner is hereby authorized to charge any additional fees or credit any overpayment to Deposit Account No. **14-1138**.  
**A duplicate copy of this sheet is enclosed.**

Date: August 23, 1999

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I hereby certify that this correspondence is being deposited with the United States Postal Service as first class mail in an envelope addressed to: Assistant Commissioner for Patents, Washington, D.C. 20231. on the date below.	
8/23/99 Date	<u>Wendy L. Harrold</u> Wendy L. Harrold

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AMENDMENT

Assistant Commissioner for Patents  
Washington, D.C. 20231  
**Box: Non-Fee Amendment**

Dear Sir:

In response to the June 8, 1999, office action, please amend the above-identified patent application as follows:

In the Claims:

Please cancel claims 44 and 81.

Please amend claims 1, 7, 13, 18, 22, 58-59, 67, and 138 as follows:

1. (Twice Amended) A method for identifying one or more of a plurality of sequences differing by one or more single-base changes, insertions, deletions, or translocations in a plurality of target nucleotide sequences comprising:
- providing a sample potentially containing one or more target nucleotide sequences with a plurality of sequence differences;